SEQUENCE LISTING





<110> Innogenetics N.V.

<120> New sequences of hepatitis C virus genotypes for diagnosis, prophylaxis and therapy

<130> 35

<160> 284

<170> PatentIn version 3.1

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<212> DNA

<213> hepatitis C virus

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213

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PRT <212>

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Ala Gln Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Pro Thr 20

Ser Phe Gly Asn Thr Ile Thr Cys Tyr Ile Lys Ala Thr Ala Ala Ala 40 35

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Ser Pho	e Gly 35	Asn	Thr	Ile	Thr	Cys 40	Tyr	Ile	Lys	Ala	Thr 45	Ala	Ala	Ala	
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Ser Ph	ne Gly 35	y Asr	Thr	: Ile	. Thr	Cys	туг	Ile	e Lys	s Ala	Thr	Ala	Ala	Ala	

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11 213

DNA

hepatitis C virus

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Cys Ser Phe 35	Ser Ile Phe	Leu Leu 40	Ala Leu	Phe Ser	Cys Leu 45	Ile His
Pro Ala Ala 50	Ser Leu Glu	Trp Arg	Asn Thr	Ser Gly	Leu Tyr	Val Leu
Thr Asn Asp	Cys Ser Asn 70	Ser Ser	Ile Val	Tyr Glu 75	Ala Asp	Asp Val
Ile Leu His	Thr Pro Gly 85	Cys Val	Pro Cys 90	Val Glr	Asp Gly	Asn Thr 95
Ser Ala Cys	Trp Thr Pro	Val Thr	Pro Thr 105	Val Ala	Val Arg 110	
Gly Ala Thr	Thr Ala Ser	: Ile Arg 120	Arg His	Val Asp	Met Leu 125	Val Gly
Ala Ala Thr 130	Met Cys Ser	Ala Leu 135	Tyr Val	Gly Asp		Gly Ala
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His Arg Met Ala 180

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<211> 180

<212> PRT

<213> hepatitis C virus

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Cys Ser Phe Ser Ile Phe Leu Pro Ala Leu Phe Ser Cys Leu Ile His 45 40

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 55

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 70

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	Ala	Ala 130	Thr	Met	Cys	Ser	Ala 135	Leu	Tyr	Val	Gly	Asp 140	Met	Cys	Gly	Ala		
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<212> PRT

<213> hepatitis C virus

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 20 25 30

Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Thr Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Ala Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val
100 105 110

Gly Ala Thr Thr Ala Ser Ile Arg Arg His Val Asp Ile Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160

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His Arg Met Ala 180

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cctctatgtc cttaccaacg actgttccaa tagcagtatt gtgtacgagg ccgatgacgt 240

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<212> PRT

<213> hepatitis C virus

<400> 20

Val Gly Ala Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val 5

Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly

Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 80 70 65

Ile Leu His Thr Pro Gly Cys Ile Pro Cys Val Gln Asp Gly Asn Thr 85

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Lys Tyr Val

100 105 110

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly 120 Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 135 Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg His Gln 150 155 Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly His Arg Met Ala 180 <210> 21 <211> . 541 <212> DNA <213> hepatitis C virus <400> 21 cgtcggcgct cccgtaggag gcgtcgcaag agcccttgcg catggcgtga gggcccttga 60 agacgggata aatttcgcaa cagggaattt gcccggttgc tccttttcta ttttccttct 120 tgctctgttc tcttgcttaa ttcatccagc agctagtcta gagtggcgga atacgtctgg 180 cctctatgtc cttaccaacg actgttccaa tagcagtatt gtgtacgagg ccgatgacgt 240 tattctgcac acacceggct gcatacettg tgtccaggac ggcaatacat ccaegtgctg 300 gaccccagtg acacctacag tggcagtcaa gtacgtcgga gcaaccaccg cttcgatacg 360 cagtcatgtg gacctattag tgggcgcggc cacgatgtgc tctgcgctct acgtgggtga 420

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<213> hepatitis C virus

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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 135 Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 155 145 Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly His Arg Met Ala 180 <210> 25 <211> 541 <212> DNA <213> hepatitis C virus <400> 25 cgtcggcgct cccgtaggag gcgtcgcaag agcccttgcg catggcgtga gggcccttga 60 qqacqqqata aacttcqcaa cagggaattt gcccggttgc tctttttcta tcttccttct 120 tgctctgttc tcttgcttaa tccatccagc agctggtcta gagtggcgga atacgtctgg 180 cctctatgtc cttaccaacg actgttccaa tagtagtatt gtgtatgagg ccgatgacgt 240 tattctgcac gcgcccggct gtgtaccttg tgtccaggac ggcaatacgt ctacatgctg 300 gaccccagta acacctacag tggcagtcag gtacgtcggg gcaaccaccg cttcgatacg 360 cagtcatgtg gacctgttag taggcgcggc cacgatgtgc tctgcgcttt acgtgggtga 420 tatgtgtggg gccgtcttcc tcgtgggaca agccttcacg ttcagacccc gccgccatca 480 aacggtccag acctgtaact gctcgctgta cccaggccat ctttcaggac atcgaatggc 540 541 t <210> 26 <211> 180 <212> PRT hepatitis C virus <213> <400> 26 Val Gly Ala Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val

Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly

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•																			
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Pro	Ala 50	Ala	Gly	Leu	Glu	Trp 55	Arg	Asn	Thr	Ser	Gly 60	Leu	Tyr	Val	Leu			.•	
Thr 65	Asn	Asp	Cys	Ser	Asn 70	Ser	Ser	Ile	Val	Tyr 75	Glu	Ala	Asp	Asp	Val 80				
Ile	Leu	His	Ala	Pro 85	Gly	Cys	Val	Pro	Cys 90	Val	Gln	Asp	Gly	Asn 95					
Ser	Thr	Cys	Trp	Thr	Pro	Val	Thr	Pro 105	Thr	Val	Ala	Val	Arg 110	Tyr	Val				
Gly	Ala	Thr 115	Thr	Ala	Ser	Ile	Arg 120	Ser	His	Val	Asp	Leu 125	Leu	Val	Gly				
Ala	Ala 130	Thr	Met	Cys	Ser	Ala 135	Leu	Tyr	Val	Gly	Asp 140	Met	Cys	Gly	Ala				
Val 145	Phe	Leu	Val	Gly	Gln 150	Ala	Phe	Thr	Phe	Arg 155	Pro	Arg	Arg	His	Gln 160				
Thr	Val	Gln	Ţhr	Cys 165	Asn	Cys	Ser	Leu	Tyr 170		Gly	His	Leu	Ser. 175	Gly	~			
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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160

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ctcg	cgcg	gca ç	gctc	ctcc	و حد	aagt	:tggg	g acc	gagat	gtg	gaag	tgto	tc g	gtace	gctta		180
agco	aaca	ict a	catg	gaco	t ac	gcct	ctto	c tat	atco	gtt	9999	cctg	jtc d	caaaa	tgaaa		240
tatg	ctt	jac a	cacc	ccat	c ac	aaaa	ataca	a tca	tggo	atg	cato	tca					287
<210 <211 <212 <213	.> 9 !> E	0 95 PRT nepat	itis	s C v	virus	3											
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Asp 1	Phe	Trp	Glu	Ser 5	Val	Phe	Thr	Gly	Leu 10	Thr	His	Ile	Asp	Ala 15	His		
Phe	Leu	Ser	Gln 20	Thr	Lys	Gln	Gln	Gly 25	Leu	Asn	Phe	Ser	Phe 30	Leu	Thr	٠.	
Ala	Tyr	Gln 35	Ala	Thr	Val	Cys	Ala 40	Arg	Ala	Gln	Ala	Pro 45	Pro	Pro	Ser		
Trp	Asp 50	Glu	Met	Trp	Lys	Cys 55	Leu	Val	Arg	Leu	Lys 60	Pro	Thr	Leu	His	· ·	
Gly 65	Pro	Thr	Pro	Leu	Leu 70	Tyr	Arg	Leu	Gly	, Pro 75	Val	Gln	Asn	Glu	Ile 80		

Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser 85 90 95

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ctga	atct	gga a	igtaa	accad	cc ag	gcaco	ctggg	g tgt	tgct	tgg	aggg	ggtc	ctc q	gcggd	cctag
cggc	cta	ctg (ttgt	cagt	c gg	gctgo	gttg	g taa	atcgt	ggg	tcat	tatco	gag (ctggg	gggca
agco	ggc	act o	gtto	caga	ac aa	aggag	ggtgt	tgt	tatca	aaca	gtad	gate	gag a	atgga	aggagt
gcto	gca	agc o	egece	cata	ac at	cgaa	acaag	gcto	caggt	aat	agco	ccaco	cag (ttcaa	aggaga
aaat	cct	tgg a	actgo	etgea	ag co	gaged	cacco	aac	caaca	aagc	tgto	catto	gag (cccgt	aatag
ctto	caa	ctg q	gcaaa	agct	ct ga	aaaco	cttct	ggo	cacaa	agca	t				
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Cys	Met	Ser	Ala 20	Asp	Leu	Glu	Val	Thr 25	Thr	Ser	Thr	Trp	Val 30	Leu	Leu
Gly	Gly	Val 35	Leu	Ala	Ala	Leu	Ala 40	Ala	Tyr	Cys	Leu	Ser 45	Val	Gly	Cys
Val	Val 50	Ile	Val	Gly	His	Ile 55	Glu	Leu	Gly	Gly	Lys 60	Pro	Ala	Leu	Val
Pro 65	Asp	Lys	Glu	Val	Leu 70	Tyr	Gln	Gln	Tyr	Asp 75	Glu	Met	Glu	Glu	Cys 80
Ser	Gln	Ala	Ala	Pro 85	Tyr	Ile	Glu	Gln	Ala 90	Gln	Val	Ile	Ala	His 95	Gln
Phe	Lys	Glu	Lys	Ile	Leu	Gly	Leu	Leu 105	Gln	Arg	Ala	Thr	Gln	Gln	Gln

Ala Val Ile Glu Pro Val Ile Ala Ser Asn Trp Gln Lys Leu Glu Thr

115 120 125

Phe Trp His Lys His
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<400> 33
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cggcctactg cttgtcagtc ggctgcgttg taatcgtggg tcatatcgag ctgggggca
agccggcact cgttccagac aaggaggtgt tgtatcaaca gtacgatgag atggaggagt
gctcgcaagc cgccccatac atcgaacaag ctcaggtaat agcccaccag ttcaaggaga
aaatccttgg actgctgcag cgagccaccc aacaacaagc tgtcattgag cccgtaatag
cttccaactg gcaaaagctt gaaaccttct ggcacaagca t
<210> 34 <211> 133 <212> PRT <213> hepatitis C virus
<400> 34
Gln Asn Glu Ile Cys Leu Thr His Pro Val Thr Lys Tyr Ile Met Ala 1 5 10 15
Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu 20 25 30
Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys 65 70 75 80

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Leu Val

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

Phe	Lys	Glu	Lys 100	Ile	Leu	Gly	Leu	Leu 105	Gln	Arg	Ala	Thr	Gln 110	Gln	Gln	
Ala	Val	Ile 115	Glu	Pro	Val	Ile	Ala 120	Ser	Asn	Trp	Gln	Lys 125	Leu	Glu	Thr	
Phe	Trp 130	His	Lys	His												
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cggc	ctad	ctg d	ettgt	cagt	c gg	gttgt	gttg	, tga	ittgt	999	tcat	atcg	ag d	ctggg	gggca	180
agco	:ggca	aat o	gtto	caga	c aa	agag	gtgt	tgt:	atca	aca	atac	gatg	ag a	atgga	agagt	240
gcto	acaa	agc t	gccc	cata	t at	cgaa	caag	cto	aggt	aat	ágct	cacc	ag t	tcaa	ıgggaa	300
aagt	cctt	gg a	attgo	tgca	g cg	gagco	acco	aac	aaca	agc	tgto	attg	ag d	ccat	agtaa	360
ctac	caad	etg g	gcaaa	agct	t ga	aggco	tttt	ggc	acaa	gca	t				•	401
<210 <211 <212 <213	.> : !> !	36 133 PRT nepat	citis	; C v	irus					٠						
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Gln 1	Asn	Glu	Ile	Cys 5	Leu	Thr	His	Pro	Ile 10	Thr	Lys	Tyr	Ile	Met 15	Ala	
Cys	Met	Ser	Ala 20	Asp	Leu	Glu	Val	Thr 25	Thr	Ser	Thr	Trp	Val 30	Leu	Leu	

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50 55 60

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Gly Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln 105 100 Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 120 115 Phe Trp His Lys His 130 <210> 37 <211> 401 <212> DNA hepatitis C virus <213> <400> 37 tccaaaatga aatctgcttg acacacccca tcacaaaata catcatggca tgcatgtcag 60 ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggggtcctc gcggccctag 120 cggcctactg cttgtcagtc ggttgtgttg tgattgtggg tcatatcgag ctggggggca 180 agccggcaat cgttccagac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240 gctcacaagc tgccccatat atcgaacaag ctcaggtgat agctcaccag ttcaaggaaa 300 aagtccttgg attgctgcag cgagccaccc aacaacaagc tgtcattgag cccatagtaa 360 401 ctaccaactg gcaaáagctt gaggcctttt ggcacaagca t <210> 38 <211> 133 <212> PRT <213> hepatitis C virus <400> Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala 10 5

Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 40 Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50 55 Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Glu Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln 100. Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 120 115 Phe Trp His Lys His 130 <210> 39 <211> 401 <212> DNA <213> hepatitis C virus <400> tccaaaatga aatctgcttg acacaccca tcacaaaata catcatggca tgcatgtcag 60 ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggggtcctc gcggccctag 120 concetacto ettotcaque ogttototto toattotogo teatategag etgggggea 180 aqccqqcaat cqttccaqac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240 qctcacaaqc tqccccatat atcgaacaag ctcaggtaat agctcaccag ttcaaggaaa 300 aagteettgg attgetgeag egageeacce aacaacaage tgteattgag eccatagtaa 360

<210> 40

<211> 133

<212> PRT

<213> hepatitis C virus

<400> 40

Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala

401

ctaccaactg gcaaaagctt gaggcctttt ggcacaagca t

15

Cys	Met	Ser	Ala	Asp	Leu	Glu	Val	Thr	Thr	Ser	Thr	Trp	Val	Leu	Leu
			20					25					30		

10

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50 55 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys 70 75 80

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

Phe Lys Glu Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln 100 105 110

Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 115 120 125

Phe Trp His Lys His

<210> 41

<211> 509

<212> DNA

<213> hepatitis C virus

<400> 41

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<210> 42

<211> 169

<212> PRT

<213> hepatitis C virus

<400> 42

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu 165

<210> 43

<211> 509

<212> DNA

<213> hepatitis C virus

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ggggccctag	gatgggtgtg	cgcgcgactc	ggaagacttc	ggaacggtcg	caaccccgtg	180
gacggcgtca	gcctattccc	aaggcgcgcc	agcccacggg	ccggtcctgg	ggtcaacccg	240
ggtacccttg	gcccctttac	gccaatgagg	gcctcgggtg	ggcagggtgg	ctgctctccc	300
ctcgaggctc	teggeetaat	tggggcccca	atgacccccg	gcgaaaatcg	cgtaatttgg	360
gtaaggtcat	cgatacccta	acgtgcggat	tcgccgatct	catggggtat	atcccgctcg	420
taggcggccc	cattgggggc	gtcgcaaggg	ctctcgcaca	cggtgtgagg	gtccttgagg	480
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<210> 44

<211> 169

<212> PRT

<213> hepatitis C virus

<400> 44

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu 165

<210> 45

<211> 580

<212> DNA

<213> hepatitis C virus

<400> 45

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<210> 46

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 46

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

																·		
Glu	Asp	Gly	Val	Asn	Tyr	Ala	Thr	Gly	Asn	Leu	Pro	Gly	Cys	Ser	Phe			
	-	35					40					45						
Ser		Phe	Ile	Leu	Ala		Leu	Ser	Cys	Leu	Thr	Val	Pro	Ala	Ser			
	50					55					60							
							_		~ 7 -	m	77.	77-7	mb	7 ~~~	200			
Ala 65	Val	Pro	Tyr	Arg	Asn 70	Ala	Ser	GIY	TTE	75	HIS	val.	THE	ASII	80			
0.5																		
Cvs	Pro	Asn	Ser	Ser	Ile	Val	Tvr	Glu	Ala	Asp	Asn	Leu	Ile	Leu	His			
-1				85			-		90					95				
Ala	Pro	Gly		Val	Pro	Cys	Val	Met	Thr	Gly	Asn	Val	Ser	Arg	Cys			
			100					105					110				_	
			_	,			_		.	D	0	T 0	a 1	777	Wa I			
Trp	Val	Gln 115	Ile	Thr	Pro	Thr	Leu 120	ser	Ala	Pro	ser	125	GIY	Ala	Vai			
Thr	Δla	Pro	Leu	Ara	Ara	Ala	Val	Asp	Tyr	Leu	Ala	Gly	Gly	Ala	Ala			
	130			5	3	135		_	-		140							
		Ser	Ala	Leu		Val	Gly	Asp	Ala			Ala	Leu	Phe	Leu 160			
145					150					155					100			
				1	— 1	m	7	D	7		ui a	717	Thr	· Val	Gln			
Val	GIY	Gin	Met	Pne 165		Tyr	Arg	PLO	170	GIII	. nis	Ala	1111	175	Gln			
Asn	Cys	Asn	Cys	Ser	· Ile	Tyr	Ser	Gly	His	Val	Thr	Gly	His	Arg	Met	·		
	-		180					185					190)				
										,				-				
Ala	ı																	
-01	10-	47												•		·		
		47 580																
		DNA hepa	,+i+i	is C	viru	ıs	,											
					V 1.1. U													
<40	00>	47	++	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	ato t	rato	iaaat	a ta	itaa.	acto	: ata	aacc	acc	ccat	tggggg	60		
cgt	cgca	aagg	gct	ctcgc	cac a	cggt	gtga	ig gg	tcct	tgag	ggad	gggg	jtaa	acta	atgcaac	120		
agg	ggaat	tta	cccg	ggttg	gct c	tttc	ctcta	at ct	ttat	tctt	gct	ctto	tct	cgts	gtctgac	180		
															-			

cgttccggcc	tctgcagttc	cctaccgaaa	tgcctctggg	atttatcatg	ttaccaatga	240
ttgcccaaac	tcttccatag	tctatgaggc	agataacctg	atcctacacg	cacctggttg	300
cgtgccttgt	gtcatgacag	gtaatgtgag	tagatgctgg	gtccaaatta	cccctacact	360
gtcagccccg	agcctcggag	cagtcacggc	tactattagg	agagccgttg	actacctagc	420
gggaggggct	gccctctgct	ccgcgttata	cgtaggagac	gcgtgtgggg	cactattctt	480
ggtaggccaa	atgttcacct	ataggcctcg	ccagcacgct	acggtgcaga	actgcaactg	540
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<210> 48

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 48

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys
100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val 115 120 125

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala 130 135 140 Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 170 175

Asn Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met 180 185 190

Ala

<210> 49

<211> 959

<212> DNA

<213> hepatitis C virus

<400> 49

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<211> 319

<212> PRT

<213> hepatitis C virus

<400> 50

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 185 190

Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205 Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 210 215 220

Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 225 230 235 240

Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 255

Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 265 270

Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 275 280 285

Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln Asn Cys 290 295 300

Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met Ala 305 310 315

<210> 51

<211> 959

<212> DNA

<213> hepatitis C virus

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Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60
Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly65707580
Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95
Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110
Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp

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	Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175	•
	Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 185 190	
	Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205	
	Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 210 215 220	
	Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 225 230 235 240	
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	Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 265 270	
	Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 275 280 285	
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Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 185 190

Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205

Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 210 215 220

Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 225 230 235 240

Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 255

Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 265 270

Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 275 280 285

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attcttggga taggcactgt ccttgaccag gcagagacgg ctggagctag gctcgtcgtc 180

ttggccacgg ncacccctcc cggcagtgtg acaacgcccc accccaacat cgaggaagtg 240
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Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr Val Leu 35 40 45

Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala Thr Xaa 50 55 60

Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu Glu Val 65 70 75 80

Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Gly Arg Ala Ile Pro

Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser Lys
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Lys Asn Cys Asp Glu Leu
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Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr Val Leu 35 40 45

Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Xaa Thr Xaa 50 55 60

Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu Glu Val 65 70 75 80

Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Xaa Arg Ala Ile Pro 85 90 95

Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser Lys
100 105 110

Lys Lys Cys Asp Glu Leu Arg Gln Ala Thr Asp Gln Pro Gly Arg Glu 115 120 125

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                                                                       180
gatgagatgg aggagtgete ggeetegttg ecetatatgg acgaaacaeg tnecattgee
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Ile Ile Xaa Asp Arg Glu Val Leu Tyr Gln Gln Phe Asp Glu Met Glu
                      55
Glu Cys Ser Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Xaa Ile Ala
                   70 -
Gly Gln Phe Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln
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Lys Ala Glu Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala
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105

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Ile Ile Pro Asp Arg Glu Ala Leu Tyr Xaa Gln Phe Asp Glu Met Glu 50 55 60

Glu Cys Ser Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala 65 70 75 80

Gly Gln Phe Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln 85 90 95

Lys Ala Glu Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala 100 105 110

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<212> PRT

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<213> hepatitis C virus
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<400> 94

Val Gln Asp Gly Asn Thr Ser Thr 1 5

<210> 95

<211> 8

<212> PRT

<213> hepatitis C virus

<400> 95

Val Gln Asp Gly Asn Thr Ser Ala
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<210> 96

<211> 10

<212> PRT

<213> hepatitis C virus

<400> 96

Val Lys Tyr Val Gly Ala Thr Thr Ala Ser 1 5 10

<210> 97

<211> 20

<212> PRT

<213> hepatitis C virus

<400> 97

Leu Gly Gly Lys Pro Ala Ile Val Pro Asp Lys Glu Val Leu Tyr Gln
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Gln Tyr Asp Glu 20

<210> 98

<211> 20

<212> PRT

<213> hepatitis C virus

<400> 98

Leu Gly Gly Lys Pro Ala Leu Val Pro Asp Lys Glu Val Leu Tyr Gln
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Gln Tyr Asp Glu

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Phe Lys Glu Lys
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Thr Gln Gln Gln
            20
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Ile Ala His Gln Phe Lys Glu Lys Ile Leu Gly Leu Leu Gln Arg Ala
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Thr Gln Gln Gln
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Leu Ser Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu Ala Leu Tyr Gln
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Gln Phe Asp Glu
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Gln Phe Asp Glu
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Phe Lys Glu Lys
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Gly Gln Lys Ala
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<400> 106

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<210> 107

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 107

Ser Thr Val Thr Glu Lys Asp Ile Arg Val Glu Glu Glu Val Tyr Gln 1 5 10 15

Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Ala Ile Thr Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Lys Ala 65 70 75 80

Ala Gly Leu Arg Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Pro Leu Gly
100 105 110

Ala

<210> 108

<211> 340

<212> DNA

<213> hepatitis C virus

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catgcataac agcaggggag acctgtgcgg gtatcgcaga tgccgtgcga gcggcgtcta 180
caccaccagc ttcgggaaca cactgacgtg ctatctcaaa gcctcagccg ctatcagagc 240
ggcggggctg agagactgca ccatgttggt ctgtggtgat gacctggtcg tcattgctga 300
aagcgatggc gtagaggagg acaaacgagc cctcggagcc 340

<210> 109

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 109

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Cys Cys Asp Leu Glu Pro Glu Ala Arg Arg Ala Ile Thr Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Arg Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ala Gly Leu Arg Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
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Ala

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caccaccagc ttcgggaaca cactgacgtg ctatctcaaa gcctcagccg ccatcagggc 240
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<210> 111

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 111

Ser Thr Val Thr Glu Lys Asp Ile Arg Val Glu Glu Glu Val Tyr Gln 1 5 10 15

Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Thr Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ser Gly Leu Arg Asp Cys Thr Met Leu Val Tyr Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
100 105 110

Val

<210> 112

<211> 340

<212> DNA

<213> hepatitis C virus

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catgcata	aac	agcaagggag	acctgtgtgg	gtatcgcaga	tgccgcgcaa	gcggcgtcta	180
caccacca	agc	ttcgggaaca	cactgacgtg	ctacctcaaa	gcctcagccg	ctatcagagc	240
ggcgggg	ctg	agagactgca	ccatgttggt	ctgtggtgat	gacctggtcg	tcatcgctga	300
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<210> 113

<211> 113

<212> PRT

<213> hepatitis C virus

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1 5 10 15

Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Ala Ile Thr Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ala Gly Leu Arg Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

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Ala

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60

120

180

240

300

340

Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Val Ala Ala Thr Arg Ala

Ala Gly Leu Lys Gly Cys Ser Met Leu Val Cys Gly Asp Asp Leu Val 85

Val Ile Cys Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg 105 100

Ala

340 <211> <212> DNA <213> hepatitis C virus <400> 116 ctccactgtg actgagagag acatcaaggt cgaagaagaa gtctatcagt gttgtgatct ggagcccgag gcccgcaagg taatagccgc cctcacggag agactctacg tgggcggccc catgcataac agcaagggag acctttgcgg gtatcgtaga tgccgcgcga gcggcgtata caccaccage ttegggaaca caatgaegtg etacettaag geeteageag eeateaggge tgcggggcta aaggattgca ccatgctggt ttgcggtgac gacctagtcg tgatcgccga gagcggtggc gttgaggagg acaaacgagc cctcggagct <210> 117 <211> 113 <212> PRT <213> hepatitis C virus <400> 117 Ser Thr Val Thr Glu Arg Asp Ile Lys Val Glu Glu Val Tyr Gln Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Ala Ala Leu Thr Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 Gly Asn Thr Met Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 75 65

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val

Val Ile Ala Glu Ser Gly Gly Val Glu Glu Asp Lys Arg Ala Leu Gly

90

110

85

100

60

120

180

240

300

340

Ala

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gttc	ccac	ct c	ggcc	gtca	a ct	atcg	caat	gcc	tcgg	gca	tcta	tcac	at (cacca	atga
tgcc	cgaa	ct c	gagc	atag	t gt	acga	gacc	gag	cacc	aca	tcct	acac	ct (cccag	ggtg
ttac	cctg	cg t	gagg	gttg	g ga	atca	gtca	cgc	tgct	ggg	tggc	cctc	ac 1	tccca	ccgt
gcgg	cgcc	tt a	catc	ggcg	c tc	cgct	tgaa	tcc	ctcc	gga	gtca	tgtg	ga	tctga	tggt
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Thr 1	Cys	Gly	Phe	Ala 5	Asp	Leu	Met	Gly	Tyr 10	Ile	Pro	Leu	Val	Gly 15	Ala
Pro	Val	Gly	Gly 20	Val	Ala	Arg	Ala	Leu 25	Ala	His	Gly	Val	Arg 30	Ala	Val
Glu	Asp	Gly 35	Ile	Asn	Tyr	Ala	Thr 40	Gly	Asn	Leu	Pro	Gly 45	Суя	Ser	Phe
Ser	Ile 50	Phe	Leu	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Thr	Ser
Ala 65	Val	Asn	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Tyr 75	His	Ile	Thi	Asn	Asp 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Thr Glu His His Ile Leu His
85 90 95

Leu Pro Gly Cys Leu Pro Cys Val Arg Val Gly Asn Gln Ser Arg Cys Trp Val Ala Leu Thr Pro Thr Val Ala Ala Pro Tyr Ile Gly Ala Pro 115 Leu Glu Ser Leu Arg Ser His Val Asp Leu Met Val Gly Ala Ala Thr 135 130 Ala Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 155 145 150 Val Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 170 165 Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg 180 <210> 120 <211> 574 <212> DNA hepatitis C virus <213> <400> 120 acgtgcggct ttgccgacct catgggatac atcccgctcg tgggcgcccc tgtgggtggc 60 gtcgccaggg ccttggcaca tggtgtcagg gccgtggagg acgggattaa ctatgcaaca 120 gggaatette eeggttgete ettttetate tteettetag eacttetete gtgettgaet 180 gtcccggcct cggcgcagca ctaccggaac atctcgggca tttatcacgt caccaatgac 240 tgcccgaact ctagtatagt gtatgaagct gaccatcata tcatgcatct accagggtgt 300 gtgccttgcg tgagaaccgg gaacacctcg cgctgctggg ttcctttaac acccactgtg 360 gctgccccct atgttggcgc gccgctcgaa tccatgcggc ggcacgtgga cttaatggtg 420 ggtgccgcca ccgtctgctc ggccctgtac atcggagacc tttgcggagg tgtcttcctg 480 gtcgggcaga tgttcacctt ccggccgcgc cgccattgga ctacccagga ctgcaactgc 540 574 tctatctatg atggccacat caccggccat agaa <210> 121

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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Gln His Tyr Arg Asn Ile Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Thr Ser Arg Cys
100 105 110

Trp Val Pro Leu Thr Pro Thr Val Ala Ala Pro Tyr Val Gly Ala Pro 115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Val Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Arg Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Asp Gly His Ile Thr Gly His Arg 180 185 190

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gggaatcttc	ccggttgctc	tttctctatc	ttcctcctgg	cacttctttc	gtgcctcact	180
gtcccagcgt	cagctgagca	ctaccggaat	gcttcgggca	tctatcacat	caccaatgac	240
tgtccgaatt	ccagcgtagt	ctatgaaact	gaccaccata	tattgcactt	gccggggtgc	300
gtaccctgcg	tgagggccgg	gaacgtgtct	cgttgctgga	cgccggtaac	acctacggtg	360
gctgccgtat	ccatggacgc	teegetegaġ	tccttccggc	ggcatgtgga	cctaatggta	420.
ggtgcggcca	ccgtgtgttc	tgtcctctat	gttggagacc	tctgtggagg	tgctttccta	480
gtggggcaga	tgttcacctt	ccagccgcgt	cgccactgga	ccacgcagga	ttgtaattgc	540
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<211> 191

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<400> 123

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Glu His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Val Val Tyr Glu Thr Asp His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Ala Gly Asn Val Ser Arg Cys
100 105 110

Trp Thr Pro Val Thr Pro Thr Val Ala Ala Val Ser Met Asp Ala Pro 115 120 125

Leu Glu		Phe	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr	
Val Cys	Ser	Val	Leu	Tyr 150	Val	Gly	Asp	Leu	Cys 155	Gly	Gly	Ala	Phe	Leu 160	
Val Gly	, Gln	Met	Phe 165	Thr	Phe	Gln	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln	
Asp Cys	s Asn	Cys 180	Ser	Ile	Tyr	Thr	Gly 185	His	Ile	Thr	Gly	His 190	Arg		
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<213> hepatitis C virus

Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile

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Glu His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile
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Val Tyr Glu Thr Glu His His Ile Leu His Leu
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Val Tyr Glu Ala Asp His His Ile Met His Leu
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Arg Pro Arg Arg His Trp Thr Thr Gln Asp
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- Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30
- Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45
- Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60
- Ile Pro Lys Asp Arg Arg Pro Thr Gly Lys Ser Trp Gly Lys Pro Gly 65 70 75 80
- Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95
- Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro 100 105 110
- Arg His Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
 115 120 125
- Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val 130 135 140
- Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
 145 150 155 160
- Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175
- Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Val Pro Val Ser Gly Leu 180 185 190
- Gln Val Lys Asn Thr Ser Ser Ser Tyr Met Val Thr Asn Asp Cys Gln 195 200 205
- Asn Ser Ser Ile Val Trp Gln Leu Arg Asp Ala Val Leu His Val Pro 210 215 220
- Gly Cys Val Pro Cys Glu Glu Lys Gly Asn Ile Ser Arg Cys Trp Ile

235	240
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30

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				245					250					255	

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Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Ala Val Met Leu Ala Ser 275 280 . 285

Gln Val Phe Ile Ile Ser Pro Gln His His Lys Phe Val Gln Asp Cys 290 295 300

Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala 305 310 315

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<211> 340

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<213> hepatitis C virus

<400> 145

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<210> 146

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 146

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Ala Cys Ser Leu Pro Glu Gln Ala Arg Thr Ala Ile His Ser Leu Thr 20 25 30

Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Leu :	Asn	Ser	Lys 45	Gly	Gln	Thr	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Phe 60	Thr	Thr	Ser	Met	
Gly 65	Asn	Thr	Ile	Thr	Cys 70	Tyr	Val	Lys	Ala	Gln 75	Ala	Ala	Cys	Lys	Ala 80	·
Ala	Gly	Ile	Ile	Ala 85	Pro	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
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Ala																
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ggc	gcca	acc	catc	ссса	gg 9	cgcg	ccga	a co	gagg	gcag	gtc	ctgg	gct	cago	ccgggt	240
acc	cttg	gcc	ccta	tatg	gg a	atga	gggc	t go	gggt	gggc	agg	gtgg	ctc	ctgt	ccccgc	300
gcg	gctc	tcg	cccg	tcgt	gg g	gccc	aaat	g ac	cccc	ggcg	cag	ga				345
	1>	148 115 PRT hepa	ıtiti	.s C	viru	ıs					<i>:</i>					
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Met 1	: Ser	Thr	Leu	pro 5	Lys	Pro	Glr	ı Arg	Lys 10	: Thr	. Lys	arg	J Ası	Thr 15	Asn .	
Pro	Gly	/ His	Arç 20	Thi	. Le	ı Ser	Ser	Glr 25	n Ala	a Ala	a Val	. Arg	30	. Leu	ı Val	

Glu Phe Thr Cys Tyr His Ala Gly Ala Pro Ser Trp Val Cys Val Gln Cys Ala Arg Leu Pro Ser Gly Arg Asn Leu Ala Val Gly Ala Asn Pro 55 Ser Pro Gly Arg Ala Glu Pro Arg Ala Gly Pro Gly Leu Ser Pro Gly 70 Thr Leu Gly Pro Tyr Met Gly Met Arg Ala Ala Gly Gly Gln Gly Gly 90 85 Ser Cys Pro Arg Ala Ala Leu Ala Arg Arg Gly Ala Gln Met Thr Pro 105 Gly Ala Gly 115 <210> 149 <211> 280 <212> DNA <213> hepatitis C virus <400> 149 ggcctgtgac ctcaaggacg aggctaggag ggtgataact tcactcacgg agcggcttta 60 ctgtggtggt cctatgttca acagcaaggg acaacactgc ggttaccgcc gctgccgtgc. 120 tagtggggtg ctacccacca gcttcgggaa cacaatcacc tgttacatca aagcaaaggc 180 agctaccaaa gctgccggaa ttaaaaatcc atcattcctt gtctgcggag atgacttggt 240 280 cgtgattgct gagagtgcag ggatcgatga ggacagagcg <210> 150 <211> 93 <212> PRT <213> hepatitis C virus 150 <400> Ala Cys Asp Leu Lys Asp Glu Ala Arg Arg Val Ile Thr Ser Leu Thr 10 Glu Arg Leu Tyr Cys Gly Gly Pro Met Phe Asn Ser Lys Gly Gln His

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Pro Thr Ser Phe 40 Gly Asn Thr Ile Thr Cys Tyr Ile Lys Ala Lys Ala Ala Thr Lys Ala 50 Ala Gly Ile Lys Asn Pro Ser Phe Leu Val Cys Gly Asp Asp Leu Val 75 Val Ile Ala Glu Ser Ala Gly Ile Asp Glu Asp Arg Ala <210> 151 <211> 499 <212> DNA hepatitis C virus <213> <400> atgagcacga atcctaaacc tcaaagaaaa accaaaagaa acaccaaccg tcgcccacag 60 gacgtcaagt tcccgggcgg tggtcagatc gttggcggag tttacttgtt gccgcgcagg 120 ggccctagga tgggtgtgcg cgcgactcgg aagacttcgg aacggtcgca accccgtgga 180 cggcgtcagc ctattcccaa ggcgcgccag cccacgggcc ggtcctgggg tcaacccggg 240 taccettgge ccetttacge caatgaggge ctcgggtggg cagggtgget geteteceet 300 cgaggetete ggeetaattg gggeeceaat gaeeceegge gaaaategeg taatttgggt 360 aaggtcatcg ataccctaac gtgcggattc gccgatctca tggggtatat cccgctcgta 420 ggeggeeeca ttgggggegt egeaaggget etegeaeaeg gtgtgagggt eettgaggae 480 499 ggggtaaact atgcaacag 152 <210> <211> 166 <212> PRT <213> hepatitis C virus <400> 152 Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala

Thr	Arg 50	Lys	Thr	Ser	Glu	Arg 55	Ser	Gln	Pro	Arg	Gly 60	Arg	Arg	Gln	Pro	
Ile 65	Pro	Lys	Ala	Arg	Gln 70	Pro	Thr	Gly	Arg	Ser 75	Trp	Gly	Gln	Pro	Gly 80	
Tyr	Pro	Trp	Pro	Leu 85	Tyr	Ala	Asn	Glu	Gly 90	Leu	Gly	Trp	Ala	Gly 95	Trp	
Leu	Leu	Ser	Pro 100	Arg	Gly	Ser	Arg	Pro 105	Asn	Trp	Gly	Pro	Asn 110	Asp	Pro	
Arg	Arg	Lys 115	Ser	Arg	Asn	Leu	Gly 120	Lys	Val	Ile	Asp	Thr 125	Leu	Thr	Cys	
Gly	Phe 130	Ala	Asp	Leu	Met	Gly 135	Tyr	Ile	Pro	Leu	Val 140	Gly	Gly	Pro	Ile	
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Gly	Val	Asn	Tyr	Ala 165			·									
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															ggttgc	300
															acactg	360
tca	gccc	cga	gcct	cgga	igc a	Igtca	cggo	ct cc	tctt	.cgga	gag	ccgt	tga	ctac	:ctagcg	420

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<210> 154

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 154

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Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 25

Glu Asp Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe 40 35

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 55 50

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 75 70 65

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 85

Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val 120 115

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala 135 130

Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 150 145

Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 175 170 165

Asn Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met

180 185 190

Ala

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gtcccggcct	ctgcagttcc	ctaccgaaat	gcctctggga	tctatcatgt	caccaatgat	240
tgcccaaact	cttccatagt	ctatgaggca	gatgatctga	tcctacacgc	acctggctgc	300
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tcagccccga	gcttcggagc	agtcacggct	ccccttcgga	gagccgttga	ttacttggtg	420
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gtaggccaaa	tgttcaccta	taggcctcgc	cagcatgcta	cggtgcagga	ctgcaactgt	540
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<211> 193

<212> PRT

<213> hepatitis C virus

<400> 156

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Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 70 Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu His 85 Ala Pro Gly Cys Val Pro Cys Val Arg Lys Asp Asn Val Ser Arg Cys Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Phe Gly Ala Val Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Val Gly Gly Ala Ala Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 155 145 150 Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 170 Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Gln Met 185 180 Ala <210> 157 <211> 530 <212> DNA hepatitis C virus <213> <400> 157 cacctacgac agetetgetg gtggcccagt tactgcggat tececaagtg gteattgaca 60 tcatcgcagg gagccactgg ggggtcttgt ttgccgccgc atactatgca tcggtggcta 120 actggaccaa ggtcgtgctg gtcttgtttc tgtttgcagg ggttgatgct actacccaga 180 tttcgggcgg ctccagcgcc caaacgacgt atggcatcgc ctcatttatc acccgcggcg 240 cgcagcagaa actgcagctc ataaatacca acggaagctg gcacatcaac aggaccgccc 300 ttaattgtaa tgacagcctc cagactgggt tcatagccgg cctcttctac taccataagt 360 tcaactcttc tggatgcccg gatcggatgg ctagctgtag ggcccttgcc acttttgacc 420 agggctgggg aactatcagc tatgccaaca tatcgggtcc cagtgatgac aaaccatatt 480 <210> 158

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<212> PRT

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Val Ile Asp Ile Ile Ala Gly Ser His Trp Gly Val Leu Phe Ala Ala 20 25 30

Ala Tyr Tyr Ala Ser Val Ala Asn Trp Thr Lys Val Val Leu Val Leu 35 40 45

Phe Leu Phe Ala Gly Val Asp Ala Thr Thr Gln Ile Ser Gly Gly Ser 50 55 60

Ser Ala Gln Thr Thr Tyr Gly Ile Ala Ser Phe Ile Thr Arg Gly Ala 65 70 75 80

Gln Gln Lys Leu Gln Leu Ile Asn Thr Asn Gly Ser Trp His Ile Asn 85 90 95

Arg Thr Ala Leu Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Ile Ala 100 105 110

Gly Leu Phe Tyr Tyr His Lys Phe Asn Ser Ser Gly Cys Pro Asp Arg 115 120 125

Met Ala Ser Cys Arg Ala Leu Ala Thr Phe Asp Gln Gly Trp Gly Thr 130 135 140

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Ala Ile Cys Glu Ser Gln Gly Thr His Glu Asp Glu Ala Ser Leu Arg 100 105 110 60

120

180

240

300

340

Ala

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caccaccagt atgggcaaca ccatgacgtg ctacattaag gctttagcct cctgtagaac
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Gln Arg Leu Phe Cys Gly Gly Pro Met Tyr Asn Ser Lys Gly Gln Gln 35 40 45
Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Met 50 55 60
Gly Asn Thr Met Thr Cys Tyr Ile Lys Ala Leu Ala Ser Cys Arg Thr 65 70 75 80

Ala Gly Leu Arg Asp Tyr Thr Leu Leu Val Cys Gly Asp Asp His Val

Ala Ile Cys Glu Ser Gln Gly Thr His Glu Asp Glu Ala Asn Leu Arg

Ala

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	~ D														
<213:		AN							-						
	> h	epat	itis	C V	irus										
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acga	geae	ga a		auuo	J. J.					_			_	-	_
									~~~	a 2 a	+++>	atta	++ 0	ccaa	acada
gacg	ttaa	gt t	cccg	ggtg	g tg	geca	gate	git	ggcg	gag	LLLA	cccg	בני ב	jeege	gcagg
ggcc	ctag	gt t	gggt	gtgc	g cg	cgac	tcgg	aag	actt	cgg	agcg	gtcg	ca a	icctc	gtggg
agge	qcca	ac c	tatc	ccca	a gg	cgcg	ccga	tcc	gagg	gca	gatc	ctgg	gc g	gcago	ccggg
	_														
tato	ctta	מכ כ	aatt	tacq	αса	atga	aaac	tat	aaat	ggg	cagg	gtgg	ct o	ctgt	cccct
	0009	500			J		J J J	_	555						
	~~+~	<b>+</b> ~ ~	~~~	+ a++	~ ~	~~~~	taat	gat	ccc	aac	agag	ataa	ca d	aacc	tgggt
egeg	ggtt	LC 9	geeg		9 99	gccc	caac	gue		350	25~5		-5		-222
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aagg	tcat	cg a	tacc	ctaa	c at	gcgg	CTTC	gcc	gacc	Lua	Lygy	atac	ac	.ccgc	ttgta
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Met	> 1	64		Pro			Gln	Arg		Thr	Lys	Arg	Asn	Thr 15	Asn
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Met 1	> 1 Ser	64 Thr	Asn	Pro 5	Lys	Leu		٠.	10					15	
Met 1	> 1 Ser	64 Thr	Asn	Pro 5	Lys	Leu		Pro	10				Ile	Thr 15 Val	
Met 1	> 1 Ser	64 Thr	Asn	Pro 5	Lys	Leu		٠.	10					15	
Met 1	> 1 Ser	64 Thr	Asn	Pro 5	Lys	Leu		Pro	10				Ile	15	
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Met 1 Arg Gly	> 1 Ser Arg Val	Thr Pro Tyr 35	Asn Met 20	Pro 5 Asp	Lys Val Pro	Leu Lys Arg	Phe Arg 40	Pro 25 Gly	Gly	Gly	Gly Leu Gly	Gln Gly 45	Ile 30 Val	15 Val	Gly Ala
Met 1 Arg Gly	> 1 Ser Arg Val	Thr Pro Tyr 35	Asn Met 20	Pro 5 Asp	Lys Val Pro	Leu Lys Arg	Phe Arg 40	Pro 25 Gly	Gly	Gly	Gly	Gln Gly 45	Ile 30 Val	Val	Gly Ala
Met 1 Arg Gly	> 1 Ser Arg Val	Thr Pro Tyr 35	Asn Met 20	Pro 5 Asp	Lys Val Pro	Leu Lys Arg	Phe Arg 40	Pro 25 Gly	Gly	Gly	Gly Leu Gly	Gln Gly 45	Ile 30 Val	Val	Gly Ala
Met 1 Arg Gly	> 1 Ser Arg Val	Thr Pro Tyr 35	Asn Met 20 Leu Thr	Pro 5 Asp Leu Ser	Lys Val Pro Glu	Leu Lys Arg Arg	Phe Arg 40	Pro 25 Gly	Gly Pro	Gly Arg	Gly Leu Gly 60	Gln Gly 45 Arg	Ile 30 Val	Val Arg	Gly Ala Pro
Met 1 Arg Gly Thr	> 1 Ser Arg Val	Thr Pro Tyr 35	Asn Met 20 Leu Thr	Pro 5 Asp Leu Ser	Lys Val Pro Glu Arg	Leu Lys Arg Arg	Phe Arg 40	Pro 25 Gly	Gly Pro	Gly Arg Arg	Gly Leu Gly 60	Gln Gly 45 Arg	Ile 30 Val	Val	Gly Ala Pro
Met 1 Arg Gly	> 1 Ser Arg Val	Thr Pro Tyr 35	Asn Met 20 Leu Thr	Pro 5 Asp Leu Ser	Lys Val Pro Glu	Leu Lys Arg Arg	Phe Arg 40	Pro 25 Gly	Gly Pro	Gly Arg	Gly Leu Gly 60	Gln Gly 45 Arg	Ile 30 Val	Val Arg	Gly Ala Pro
Met 1 Arg Gly Thr	> 1 Ser Arg Val	Thr Pro Tyr 35	Asn Met 20 Leu Thr	Pro 5 Asp Leu Ser	Lys Val Pro Glu Arg	Leu Lys Arg Arg	Phe Arg 40	Pro 25 Gly	Gly Pro	Gly Arg Arg	Gly Leu Gly 60	Gln Gly 45 Arg	Ile 30 Val	Val Arg	Gly Ala Pro
Met 1 Arg Gly Thr	> 1 Ser Arg Val Arg 50	Thr Pro Tyr 35 Lys	Asn Met 20 Leu Thr	Pro 5 Asp Leu Ser	Lys Val Pro Glu Arg	Leu Lys Arg Arg 55	Phe Arg 40 Ser	Pro 25 Gly Gln	Gly Pro Pro	Gly Arg Arg Ser	Gly Leu Gly 60	Gln Gly 45 Arg	Ile 30 Val Arg	Val Arg Gln Pro	Gly Ala Pro Gly 80
Met 1 Arg Gly Thr	> 1 Ser Arg Val Arg 50	Thr Pro Tyr 35 Lys	Asn Met 20 Leu Thr	Pro 5 Asp Leu Ser	Lys Val Pro Glu Arg	Leu Lys Arg Arg 55	Phe Arg 40 Ser	Pro 25 Gly Gln	Gly Pro Pro	Gly Arg Arg Ser	Gly Leu Gly 60	Gln Gly 45 Arg	Ile 30 Val Arg	Val Arg	Gly Ala Pro Gly 80

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro

100

110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val 135 Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp 155 150 145 Gly Ile Asn Tyr Ala Thr 165 <210> 165 <211> 499 <212> DNA <213> hepatitis C virus <220> <221> misc_feature (306)..(306) <222> "n" is any nucleotide <223> <220> <221> misc feature (313)..(313) <222> "n" is any nucleotide <223> <220> <221> misc feature <222> (340)..(340) "n" is any nucleotide <223> <400> atgagcacga atcctaaacc tcaaagaaaa accaaacgta acaccaaccg ccgccctatg 60 gacgttaagt tcccaggcgg tggtcagatc gttggcggag tttacttgtt gccgcgcagg 120 ggccccaggt tgggtgtgcg cgcgactcgg aagacttcgg agcggtcgca acctcgtggg 180 aggegecaae etatececaa ggegegeega acegagggea gateetggge geageeeggg . tateettgge ceetttaegg caatgaggge tgtgggtggg cagggtgget cetgteecet 300 cgcggntctc ggncgtcttg gggccccaat gatccccggn ggagatcccg caacttgggt 360 aaggtcatcg ataccctaac atgcggcttc gccgacctca tgggatacat cccgcttgta 420

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<213> hepatitis C virus

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<223> "Xaa" is any amino acid

<220>

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<223> "Xaa" is any amino acid

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<222> (114)..(114)

<223> "Xaa" is any amino acid

<400> 166

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Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Arg Thr Glu Gly Arg Ser Trp Ala Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Xaa Ser Arg Xaa Ser Trp Gly Pro Asn Asp Pro 100 105 110

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<210> 168

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 168

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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 50 55 60

Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Ala Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Met Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln
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Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

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<211> 579

<212> DNA

<213> hepatitis C virus

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<211> 193

<212> PRT

<213> hepatitis C virus

<400> 170

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Gly Val Asn Tyr Arg Asn Ala Ser Gly Val Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Asp Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Lys Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Val Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Val Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Ile Thr Gly His Arg Met 180 185 190

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<210>	172	•				•	
<211>	193			·			
<212>	PRT						

<400> 172

<213> hepatitis C virus

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val His Tyr His Asn Thr Ser Gly Ile Tyr His Leu Thr Asn Asp

Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Phe	Glu	Ala 90	Val	His	His	Ile	Leu 95	His	
Leu	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Arg 105	Thr	Gly	Asn	Gln	Ser 110	Arg	Cys	
Trp	Val	Ala 115	Leu	Thr	Pro	Thr	Leu 120	Ala	Ala	Pro	Tyr	Leu 125	Gly	Ala	Pro	
	Glu 130	Ser	Met	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Thr	Ala	Thr	
Leu 145	Cys	Ser	Ala	Leu	Tyr 150	Val	Gly	Asp	Leu	Cys 155	Gly	Gly	Ile	Phe	Leu 160	
Ala	Gly	Gln	Met	Phe 165	Thr	Phe	Arg	Pro	Arg 170	Leu	His	Trp	Thr	Thr 175	Gln	-
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gco	gcgo	ccac	aca	ttggd	egc t	cca	cttga	ag to	ccat	gcgad	gto	catgt	gga	tttg	gatggta	420

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<210> 174

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 174

Thr Cys Gly Ser Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val His Tyr His Asn Thr Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Phe Glu Ala Glu His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Ile Ala Leu Thr Pro Thr Leu Ala Ala Pro His Ile Gly Ala Pro 115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Thr Ala Thr 130 135 140

Leu Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Ile Phe Leu 145 150 155 160

Val Gly Gln Met Phe Asn Phe Arg Pro Arg Leu His Trp Thr Thr Gln 165 170 175

Glu Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met

180 185 190

Ala

	175 579 DNA hepa	atitis C vir	rus				
	175	,					-
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gggaato	cttc	ccggttgctc	cttttctatc	ttccttctag	cacttctctc	gtgcttgact	180
gtcccg	gcct	cggcgcagca	ctaccggaac	atctcgggca	tttatcacgt	caccaatgac	240
tgcccga	aact	ctagtatagt	gtatgaagct	gaccatcata	tcatgcatct	accagggtgt	300
gtgcctt	tgcg	tgagaaccgg	gaacacctcg	cgctgctggg	ttcctttaac	acccactgtg	360
gctgcc	ccct	atgttggcgc	gccgctcgaa	tccatgcggc	ggcacgtgga	cttaatggtg	420
ggtgcc	gcca	ccgtctgctc	ggccctgtac	atcggagacc	tttgcggagg	tgtcttcctg	480
gtcggg	caga	tgttcacctt	ccggccgcgc	cgccattgga	ctacccagga	ctgcaactgc	540
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<210> 176

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 176

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala 65	Gln	His	Tyr	Arg	Asn 70	Ile	Ser	Gly	Ile	Tyr 75	His	Val	Thr	Asn	Asp 80	
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr	Glu	Ala 90	Asp	His	His	Ile	Met 95	His	
Leu	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Arg 105	Thr	Gly	Asn	Thr	Ser 110	Arg	Cys	
Trp	Val	Pro 115	Leu	Thr	Pro	Thr	Val. 120	Ala	Ala	Pro	Tyr	Val 125	Gly	Ala	Pro	
Leu	Glu 130	Ser	Met	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr	
Val 145	Cys	Ser	Ala	Leu	Tyr 150	Ile	Gly	Asp	Leu	Cys 155	Gly	Gly	Val	Phe	Leu 160	
Val	Gly	Gln	Met	Phe 165	Thr	Phe	Arg	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln	
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gtg	lccct	gtg	taag	aact	gg g	aatg	tctc	g cg	ttgc	tgga	ttc	cttt	aac	accc	actgta	36
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aa s	ıgcgg	сса	cctt	atgo	tc t	gccc	tcta	c at	cgga	gacc	att	gcgg	agg	tgtc	ttcttg	48

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<210> 178

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 178

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5. 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Val Leu Leu Ser Arg Leu Thr Val Pro Ala Ser 50 55 60

Ala Gln His Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Val Ser Arg Cys 100 105 110

Trp Ile Pro Leu Thr Pro Thr Val Ala Val Pro Tyr Leu Gly Ala Pro 115 120 125

Leu Thr Ser Val Arg Gln His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Leu Cys Ser Ala Leu Tyr Ile Gly Asp His Cys Gly Gly Val Phe Leu 145 150 155 160

Ala Gly Gln Met Val Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln
165 170 175

Asp Cys Asn Cys Ser Ile Tyr Val Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

<223>

"Xaa" is any amino acid

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<210>
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      hepatitis C virus
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<221> misc feature
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                                                                      120
gggaatette etggttgete tttetetate tecettettg aacttetete gtgeetgaet
                                                                      180
gttcccgcct cagccatcca ctatcgcaat gcttcggacg gttattatat caccaatgat
                                                                      240
tgcccgaact ctagcatagt gtatgaagcc gagaaccaca tettgcacet tecggggtgt
                                                                      300
ataccetgtg tgaagacegg gaatcagteg eggtgetggg tggeteteae eeceaegetg
                                                                      360
geggeeceae acetaegtge teegettteg teettaeggg egeatgtgga cetaatggtg
                                                                      420
ggggccgcca cggcatgctc cgctttttac attggagatc tgtgcggggg tgtgtttttg
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gegggecaac tgttcactat ceggecacge atteatgaaa ceaeteagga etgeaattge
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<400> 180

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Xaa Leu Ala His Gly Val Arg Ala Leu 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Ser Leu Leu Glu Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Ile His Tyr Arg Asn Ala Ser Asp Gly Tyr Tyr Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Ile Pro Cys Val Lys Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Leu Ala Ala Pro His Leu Arg Ala Pro 115 120 125

Leu Ser Ser Leu Arg Ala His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Ala Cys Ser Ala Phe Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Ala Gly Gln Leu Phe Thr Ile Arg Pro Arg Ile His Glu Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Ile Thr Gly Xaa Xaa Xaa 180 185 190

Xaa

<210> 181 <211> 579

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gggaatette etggttgete tttetetate tneettetgg eacttetete gtgeetgaet
                                                                      180
gteccggeet eggeteagea etaceggaat gtetegggea tetaceaegt eaceaatgat
                                                                      240
tgcccgaatt ccagcatagt gtatgaagcc gatcaccaca tcatgcactt accagggtgc
                                                                      300
ataccetgeg tgaggacegg gaacgttteg egetgetggg tatetetgae acetaetgtg
                                                                      360
gctgctccct acctcggggc tccgcttacg tcgctacggc ggcatgtgga tttgatggtg
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ggtgcagcca ccctttgctc tgccctctac gtcggagacc tctgtggagg tgtcttccta
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gtgggacaga tgttcacctt ccagccgcgc cgccactgga ccactcagga ctgcaactgc
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<213> hepatitis C virus
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<400> 182

<222>

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"Xaa" is any amino acid

<212>

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Ala Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Xaa Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser

Ala 65	Gln	His	Tyr	Arg	Asn 70	Val	Ser	Gly	Ile	Tyr 75	His	Val	Thr	Asn	Asp 80		
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr	Glu	Ala 90	Asp	His	His	Ile	Met 95	His		
Leu	Pro	Gly	Cys	Ile	Pro	Cys	Val	Arg 105	Thr	Gly	Asn	Val	Ser	Arg	Cys		
Trp	Val	Ser	Leu	Thr	Pro	Thr	Val 120	Ala	Ala	Pro	Tyr	Leu 125	Gly	Ala	Pro		
Leu	Thr 130	Ser	Leu	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr		
Leu 145	CÀR	Ser	Ala	Leu	Tyr 150	Val	Gly	Asp	Leu	Сув 155	Gly	Gly	Val	Phe	Leu 160		
Val	Gly	Gln	Met	Phe 165		Phe	Gln	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln		
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															gcaaca		120
															ctgact		18
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tac	ссаа	att	cgac	cata	iat c	taco	aggo	t ga	ctac	caca	tco	taca	cct	ccct	gggtgc	:	30

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gcggcgcctt	acgttggtgc	tccgctagaa	tccctccgga	gtcatgtgga	tctgatggta	420
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<210> 184

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 184

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 50 55 60

Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Ala Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Met Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln

165 170 175

Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met 180 185 190

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gggaato	ctcc	ccggttgctc	tttctctatc	tacctcttgg	cacttctctc	gtgcctgact	180
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ttaccct	gtg	tgagggttgg	gaatcagtca	cgttgttggg	tggccctctc	tcccaccgtg	360
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gttggt	caga	tgttttcttt	ccggccacga	cgccactgga	ctacgcagga	ctgcaattgt	540
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<210> 186

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 186

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Glu His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

	Ser Ile Tyr 50	Leu Leu Ala	Leu Leu Ser 55		Thr Val Pro T 50	Thr Ser	
	Ala Ile His 65	Tyr Arg Asn 70	Ala Ser Gly	Val Tyr I	His Val Thr A	Asn Asp 80	
	Cys Pro Asn	Ser Ser Ile 85	Val Tyr Glu	Ala Asp 1	His His Ile I	Leu His 95	
	Leu Pro Gly	Cys Leu Pro 100	Cys Val Arg		Asn Gln Ser <i>I</i> 110	Arg Cys	
	Trp Val Ala	Leu Ser Pro	Thr Val Ala	Ala Pro'	Tyr Ile Gly <i>I</i> 125	Ala Pro	
	Val Glu Ser 130	Phe Arg Arg	His Val Asp 135	Met Met	Val Gly Ala <i>I</i> 140	Ala Thr	
	Val Cys Ser 145	Ala Leu Tyr 150	Ile Gly Asp	Leu Cys	Gly Gly Val I	Phe Leu 160	
	Val Gly Gln	Met Phe Ser 165	Phe Arg Pro	Arg Arg	His Trp Thr	Thr Gln 175	
	Asp Cys Asn	Cys Ser Ile 180	Tyr Ala Gly		Thr Gly His (Gly Met	•
	Ala					·	
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gcgg	cgcc	tt a	catc	ggcg	c tc	cgct	tgaa	tcc	ctcc	gga	gtca	tgtg	ga t	ctga	tggta	a 420
ggtg	ccgc	ta c	tgcg	tgct	c cg	ctct	ttac	atc	ggag	acc	tgtg	cggt	gg c	gtat	ttttg	480
gttg	gtca	ga t	gttc	tctt	t cc	agcc	gcgg	cgc	cact	gga	ctac	gcag	ga c	tgca	attgt	540
tcca	tcta	cg c	gggg	cacg	t ta	cggg	ccac	agg	atgg	ca						579
<210 <211 <212 <213	> 1 > P	88 93 RT epat	itis	Сv	irus											
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Thr 1	Cys	Gly	Phe	Ala 5	Asp	Leu	Met	Gly	Tyr 10	Ile	Pro	Leu	Val	Gly 15	Ala	
Pro	Val	Gly	Gly 20	Val	Ala	Arg	Ala	Leu 25	Ala	His	Gly	Val	Arg 30	Ala	Val	
Glu	Asp	Gly 35	Ile	Asn	Tyr	Ala	Thr 40	Gly	Asn	Leu	Pro	Gly 45	Cys	Ser	Phe	
Ser	Ile 50	Phe	Leu	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Thr	Ser	
Ala 65	Val	Asn	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Tyr 75	His	Ile	Thr	Asn	Asp 80	
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr	Glu	Thr 90	Glu	His	His	Ile	Leu 95	His	
Leu	Pro	Gly	Cys 100	Leu	Pro	Cys	Val	Arg 105	Val	Gly	Asn	Gln	Ser 110	Arg	Cys	
Trp	Val	Ala 115		Thr	Pro	Thr	Val 120	Ala	Ala	Pro	Tyr	Ile 125	Gly	Ala	Pro	
Leu	Glu 130	Ser	Leu	Arg	Ser	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr	
Ala 145		Ser	Ala	Leu	Tyr 150		Gly	Asp	Leu	Cys 155	Gly	Gly	Val	Phe	Leu 160	

Val Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg Met 180 185 190

Ala

<210>	189		
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<213>	hepatitis	C	virus

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<210> 190 <211> 193

<212> PRT

<213> hepatitis C virus

<400> 190

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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe

	Ile 50	Phe	Leu	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Ala	Ser			
717	Glu.	uic	TTs zac	λνα	Λan	Δla	Ser	Glv	Tle	Tvr	His	Ile	Thr	Asn	Asp			
65	GIU	птъ	TYL	Arg	70	AIG	501	O _T y	110	75					80			
_	_	_	_		**- 1	**- 7	ID	a 1		A a n	ціс	иiс	Tle	I.e.i	Wie ·			
Cys	Pro	Asn	Ser	ser 85	vaı	vaı	Tyr	GIU	90	Asp	ura	UTS	116	95	1115			
											_	1	a	•	G			
Leu	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Arg 105	Ala	GIŸ	Asn	vai	110	Arg	Cys			
													-					
Trp	Thr	Pro 115	Val	Thr	Pro	Thr	Val 120	Ala	Ala	Val	Ser	Met 125	Asp	Ala	Pro			
Leu	Glu 130	Ser	Phe	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr			•
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Val 145	Cys	Ser	Val	Leu	Tyr 150		Gly	Asp	Leu	Cys 155	Gly	Gly	Ala	Phe	Leu 160			
143					130													
Val	Gly	Gln	Met			Phe	Gln	Pro	Arg		His	Trp	Thr	Thr 175	Gln			
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Ala																		
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	0>																	
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gac	gtta	agt	tccc	gggc	gg t	ggcc	agat	c gt	tggt	ggag	g ttt	actt	gtt	gccc	gcgcagg	1,:	20	
ggc	ccca	ıggt	tggg	gtgtg	geg o	gcga	actaç	gg aa	agact	tcgg	ago	ggto	gca	acct	.cgtggg	1	80	
aga	cgto	agc	ctat	cccc	aa g	gcac	gtcg	ga to	tgag	gggaa	a ggt	cctç	iggc	tcas	jeceggg	2	40	
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Arg Arg	Pro	Met 20	Asp	Val	Lys	Phe	Pro 25	Gly	Gly	Gly	Gln	Ile 30	Val	Gly	
Gly Val	Tyr 35	Leu	Leu	Pro	Arg	Arg 40	Gly	Pro	Arg	Leu	Gly 45	Val	Arg	Ala	
Thr Arg	Lys	Thr	Ser	Glu	Arg 55	Ser	Gln	Pro	Arg	Gly 60	Arg	Arg	Gln	Pro	
Ile Pro	Lys	Ala	Arg	Arg 70	Ser	Glu	Gly	Arg	Ser 75	Trp	Ala	Gln	Pro	Gly 80	
Tyr Pro	Trp	Pro	Leu 85	Tyr	Gly	Asn	Glu	Gly 90	Cys	Gly	Trp	Ala	Gly 95	Trp	
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ggcccc															180
aggcgto	•														240
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aaggtca	atcg	atac	cctg	ac g	tgcg	gctt	c gc	cgac	ctca	tgg	gata	cat	cccg	ctcgtg	420
ggcgcc	ccg	tcgg	gggc	gt c	gcca	gggc	c ct	ggcg	catg	gcg	tcag	ggc	tgtg	gaggac	480
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gggattaact atcgacag

<210> 194

<211> 166

<212> PRT

<213> hepatitis C virus

<400> 194

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Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Ala Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro 100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp 145 150 155 160

Gly Ile Asn Tyr Arg Gln 165

<210> 195

<211> 579

<212> DNA

<213> hepatitis C virus

<400> 195

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gggaatctgc	ctggttgctc	tttctctatc	ttcattcttg	cacttctctc	gtgcctcact	180
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tgcccaaact	cttccatagt	ctatgaggca	gatgatctga	tcctacacgc	acctggctgc	300
gtgccttgtg	tcaggaaaga	taatgtgagt	aġgtgctggg	tccaaattac	ccccacgctg	360
tcagccccga	gcttcggagc	agtcacggct	ccccttcgga	gagccgttga	ttacttggtg	420
ggaggggctg	ccctctgctc	cgcgttatac	gttggagacg	cgtgtggggc	actatttttg	480
gtaggccaaa	tgttcaccta	taggcctcgc	cagcatgcta	cggtgcagga	ctgcaactgt	540
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<210> 196

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 196

Thr Cys Gly Phe Ala Asp Leu Val Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Arg Lys Asp Asn Val Ser Arg Cys
100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Phe Gly Ala Val

115

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Val Gly Gly Ala Ala 135 130 Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 155 150 Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 170 165 Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Gln Met 185 Ala <210> 197 1485 <211> <212> DNA <213> hepatitis C virus <220> <221> misc feature <222> (787)..(787) "n" is any nucleotide <223> <220> <221> misc feature <222> (1011) . . (1011) <223> "n" is any nucleotide <220> <221> misc feature <222> (1444)..(1444) <223> "n" is any nucleotide <400> 197 tgtgccagga ccatcaccac cggagcttct atcacatact ccacttacgg caagttcctt 60 gctgatggag ggtgttcagg cggcgcgcat gacgtgatca tatgcgacga gtgccattcc 120 caggacgcca ccaccattct tgggataggc actgtccttg accaggcaga gacggctgga getaggeteg tegtettgge caeggecaee ceteceggea gtgtgacaae geeceaeeee 240 aacatcgagg aagtggccct gcctcaggag ggggaggttc ccttctacgg cagagccatt

ccccttgctt ttataaaggg tggtaggcat ctcatcttct gccattccaa gaaaaaatgt 360 gatgaactcg ccaagcaact gaccagcctg ggcgtgaacg ccgtggcata ttatagaggt 420 ctagacgtcg ccgtcatacc cacaacagga gacgtggtcg tgtgcagcac cgacgcgctc 480 atgacgggat tcaccggcga ctttgattct gtcatagact gcaactccgc cgtcactcag 540 acggtggact tcagtctgga tcccactttt accattgaga ctaccacagt gccccaggac 600 gcagtgtcca gaagccagcg ttggggccgc acggggagag gtaggcacgg catataccgg 660 tatgtctcgg ctggagagag accgtctggc atgttcgact ccgtggtgct ctgtgagtgc 720 tacgatgccg gatgtgcatg gtacgatctg actcctgccg agactaccgt gaggttgcgc 780 gcttacntaa acaccccgg gctccctgtc tgtcaggacc atttggaatt ctgggagggg 840 gtgttcacgg ggctcactaa catcgacgct cacatgctgt cacagaccaa acagggtggg 900 gagaatttcc cataccttgt agcgtaccaa gcaacagtct gtgttcgcgc gaaagcgccc 960 cccccagct gggacacaat gtggaaatgc atgctccgtc tcaaaccgac nttaactggc 1020 cctactcccc tcttgtacag gctggggccc gtccagaatg agatcacact gacgcacccc 1080 1140 atcaccaagt acattatggc ttgcatgtct gcggacttgg aggtcattac cagcacttgg gttctggtgg ggggcgttgt ggcggccctg gcggcctact gcttgacggt gggttcggta 1200 gccatagtcg gtaggatcat cctctctggg aaacctgcca tcattcccga tagggaggta 1260 ttataccagc aatttgatga gatggaggag tgctcggcct cgttgcccta tatggacgaa 1320 acacgtgcca ttgccggaca attcaaagag aaagtgctcg gcttcatcag cacgaccggc 1380 cagaaggctg aaactctgaa gccggcagcc acgtctgtgt ggaacaaggc tgagcagttc 1440 1485 tggnccacat acatgtggaa cttcatcagt gggatacaat aatag

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<210> 198
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<211> 484

<212> PRT

<213> hepatitis C virus

<220>

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<222> (263)..(263)

<223> "Xaa" is any amino acid

<220>

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<220>

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<222> (482)..(482)

<223> "Xaa" is any amino acid

<400> 198

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Gly Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala His Asp Val 20 25 30

Ile Ile Cys Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly 35 40 45

Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val 50 55 60

Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro 65 70 75 80

Asn Ile Glu Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr 85 90 95

Gly Arg Ala Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile 100 105 110

Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr 115 120 125

Ser Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala 130 135 140

Val Ile Pro Thr Thr Gly Asp Val Val Cys Ser Thr Asp Ala Leu 145 150 155 160

Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser 165 170 175

Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 180 185 190

- Glu Thr Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Trp 195 200 205
- Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala 210 215 220
- Gly Glu Arg Pro Ser Gly Met Phe Asp Ser Val Val Leu Cys Glu Cys 225 230 235 240
- Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr 245 250 255
- Val Arg Leu Arg Ala Tyr Xaa Asn Thr Pro Gly Leu Pro Val Cys Gln 260 270
- Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile 275 280 285
- Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Gly Glu Asn Phe Pro 290 295 300
- Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro 305 310 315 320
- Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro 325 330 335
- Xaa Leu Thr Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln 340 345 350
- Asn Glu Ile Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys 355 360 365
- Met Ser Ala Asp Leu Glu Val Ile Thr Ser Thr Trp Val Leu Val Gly 370 375 380
- Gly Val Val Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val 385 390 395 400
- Ala Ile Val Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro 405 410 415
- Asp Arg Glu Val Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser

425

420

Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe 435 440 Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu 455 Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe 475 470 Trp Xaa Thr Tyr <210> 199 <211> 1485 <212> DNA hepatitis C virus <213> <220> <221> misc_feature (684)..(684) <222> "n" is any nucleotide <223> <220> <221> misc feature (909)..(909) <222> <223> "n" is any nucleotide <220> <221> misc feature <222> (1058)..(1058) <223> "n" is any nucleotide <220> <221> misc_feature (1133)..(1133)<222> "n" is any nucleotide <223> <400> 199 tgtgccagga ccatcaccac cggagcttct atcacatact ccacttacgg caagttcctt 60 gctgatggag ggtgttcagg cggcgcgtat gacgtgatca tatgcgacga gtgccattcc 120 caggacgcca ccaccattct tgggataggc actgtccttg accaggcaga gacggctgga 180 getaggeteg tegtettgge caeggecaec ceteceggea gtgtgaeaac geeceaecee 240 aacatcgagg aagtggccct gcctcaggag ggggaggttc ccttctacgg cagagccatt 300 ccccttgctt ttataaaggg tggtaggcat ctcatcttct gccattccaa gaaaaaatgt 360 gatgaactcg ccaagcaact gaccagcctg ggcgtgaacg ccgtggcata ttatagaggt 420 ctagacgtcg ccgtcatccc cacagcagga gacgtggtcg tgtgcagcac cgacgcgctc 480 atgacgggat tcaccggcga ctttgattct gtcatagact gcaactccgc cgtcactcag 540 acggtggact tcagtctgga tcccactttt accattgaga ctaccacagt gccccaggac 600 gcagtgtcca gaagccagcg taggggccgc acggggagag gtaggcacgg catataccgg 660 tatgtctcgg ctggagagag accntctgac atgttcgact ccgtggtgct ctgtgagtgc 720 tacgatgccg gatgtgcgtg gtatgatctg actcctgccg agactaccgt gaggttgcgc 780 gcttacataa acaccccgg gctccctgtc tgtcaggacc atttggaatt ctgggagggg 840 gtgttcacgg ggctcactaa catcgacgct cacatgctgt cacagaccaa acagggtggg 900 gagaatttnc cataccttgt agcgtaccaa gcaacagtct gtgttcgcgc gaaagcgccc 960 cccccagct gggacacaat gtggaaatgc atgctccgtc tcaaaccgac tttaactggc 1020 cctactcccc tcttgtacag gctggggccc gtccagantg agatcacact gacgcacccc 1080 atcaccaagt acattatggc ttgcatgtct gcggacttgg aggtcattac cancacttgg 1140 gttctggtgg ggggcgttgt ggcggccctg gcggcctact gcttgacggt gggttcggta 1200 gccatagtcg gtaggatcat cctctctggg aaacctgcca tcattcccga tagggaggca 1260 ttataccagc aatttgatga gatggaggag tgctcggcct cgttgcccta tatggacgag 1320 acacgtgcca ttgccggaca attcaaagag aaagtgctcg gcttcatcag cacgaccggc 1380 cagaaggetg aaactetgaa geeggeagee aegtetgtgt ggaacaagge tgageagtte 1440 1485 tgggccacat acatgtggaa cttcatcagc gggatacaat aatag

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<210> 200
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<211> 484

<212> PRT

<213> hepatitis C virus

<220>

<221> MISC FEATURE

<222> (228)..(228)

<223> "Xaa" is any amino acid

<220>

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<222> (303)..(303)

<223> "Xaa" is any amino acid <220> <221> MISC FEATURE <222> (353)..(353) <223> "Xaa" is any amino acid <220> <221> MISC FEATURE <222> (378)..(378) <223> "Xaa" is any amino acid <400> 200 Cys Ala Arg Thr Ile Thr Thr Gly Ala Ser Ile Thr Tyr Ser Thr Tyr · 5 Gly Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Val 20 Ile Ile Cys Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr 90 Gly Arg Ala Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile 100

Phe Cys His Ser Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr

Ser Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala

Val Ile Pro Thr Ala Gly Asp Val Val Cys Ser Thr Asp Ala Leu

135

155

120

115

130

125

- Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser 165 170 175
- Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 180 185 190
- Glu Thr Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg
 195 200 205
- Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala 210 215 220
- Gly Glu Arg Xaa Ser Asp Met Phe Asp Ser Val Val Leu Cys Glu Cys 225 230 235 240
- Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr 245 250 255
- Val Arg Leu Arg Ala Tyr Ile Asn Thr Pro Gly Leu Pro Val Cys Gln 260 265 270
- Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile 275 280 285
- Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Glu Asn Xaa Pro 290 295 300
- Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro 305 310 315 320
- Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro 325 330 335
- Thr Leu Thr Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln 340 345 350
- Xaa Glu Ile Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys 355 360 365
- Met Ser Ala Asp Leu Glu Val Ile Thr Xaa Thr Trp Val Leu Val Gly 370 380
- Gly Val Val Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val

	205	400
3 90	395	400

Ala Ile Val Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro
405 410 415

Asp Arg Glu Ala Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser 420 425 430

Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe 435 440 445

Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu 450 455 460

Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe 465 470 475 480

Trp Ala Thr Tyr

<210> 201

385

<211> 340

<212> DNA

<213> hepatitis C virus

<400> 201

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tatgtacaat agcaagggag acctttgcgg gtatcgcagg tgccgcgcaa gcggcgtata 180
taccaccagc ttcgggaaca cactgacgtg ctaccttaaa gcctcagcag ccatcagggc 240
tgcggggctg aaggactgca ccatgctggt ttgcggtgac gacttagtcg tgatcgctga 300
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<210> 202

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 202

Ser Thr Val Thr Glu Arg Asp Ile Arg Val Glu Glu Glu Val Tyr Gln 1 5 10 15

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Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Tyr	Asn	Ser	Lys 45	Gly	Asp	Leu	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Tyr 60	Thr	Thr	Ser	Phe	
Gly 65	Asn	Thr	Leu	Thr	Cys 70	Tyr	Leu	Lys	Ala	Ser 75	Ala	Ala	Ile	Arg	Ala 80	
Ala	Gly	Leu	Lys	Asp 85	Cys	Thr	Met	Leu	Val 90	Сув	Gly	Asp	Asp	Leu 95	Val	
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Ala															·	
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cat	gcac	aac	agca	gggg	ag a	ccta	tgcg	g gt	accg	taga	`tgc	cgcg	cga	gcgg	cgtata	180
cac	caca	agc	ttcg	ggaa	ca c	tctg	acgt	g ct	tcct	caag	gcc	acag	cgg	ccac	caaagc	240
cgc	tggc	cta	aagg	actg	ca c	catg	ttgg	t gt	gtgg	tgac	gac	ttag	tcg	ttat	cgccga	300
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Cys Cys Asp Leu Glu Pro Glu Thr Arg Lys Val Ile Ser Ala Leu Thr Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Arg Gly Asp Leu 40 Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 55 Gly Asn Thr Leu Thr Cys Phe Leu Lys Ala Thr Ala Ala Thr Lys Ala Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Arg Arg Ala Leu Gly 105 Ala <210> 205 <211> 340 <212> DNA <213> hepatitis C virus <400> 205 ctccacggtg accgaaaggg atatcaggac cgaggaagag atctaccagt gctgcgacct 60 ggagcccgaa gcccgcaagg tgatatccgc cctaacggaa agactctacg tgggcggtcc 120 catgtacaac tccaaggggg acctatgcgg gcaacggagg tgccgcgcaa gcggggtcta 180 caccaccage ttegggaaca etgtaaegtg ttateteaag geegttgegg etaetaggge 240 cgcaggtctg aaaggttgca gcatgctggt ttgtggagac gacttagtcg tcatctgcga 300 gageggegge gtagaggagg atgeaagage ceteegagee 340 <210> 206 <211> 113 <212> hepatitis C virus <213>

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Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Tyr	Asn	Ser	Lys 45	Gly	Asp	Leu			
Cys	Gly 50	Gln	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Tyr 60	Thr	Thr	Ser	Phe			
Gly 65	Asn	Thr	Val	Thr	Cys	Tyr	Leu	Lys	Ala	Val 75	Ala	Ala	Thr	Arg	Ala 80			
Ala	Gly	Leu	Lys	Gly 85	Cys	Ser	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val			
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															cggcc		12	0
cat	gtat	aac	agca	aggg	ggg a	ccta	itgc	ia ad	etteg	gagg	tgo	cgcg	gcaa	gcgg	ggtat	a .	18	0
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tgo	agg	gctg	aaag	gatte	gca d	ctato	gctgg	gt at	gegg	gtgad	gad	ttag	gtcg	ttat	tgccg	a	30	0
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Glu Arg Leu Tyr Lys Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Leu Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala 65 70 75 80

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg 100 105 110

Ala

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tatgcacaac agcaagggag acctgtgtgg catccgtaga tgccgcgcga gcggcgttta
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caccacgage tteggaaaca egetgaettg etaceteaaa geeacagegg eeaccaggge
                                                                     240
cgcgggcttg aaggattgca ccatgctggt ctgcggngac gacctggttg tcattgctga
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gagcattggc atagacgagg acaagcaagc cctccgnact
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Cys Gly Ile Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala 65 70 75 80

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

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catqttcaac aqcaaqqqag acctgtgcgg acaacgccgg tgccgcgcaa gcggcgtgtt
caccaccage ttegggaaca cactgaegtg ctacettaaa gecacagetg ctactagage
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agcoggotta aaagattgca coatgotggt ctgcggtgac gacttagtcg ttatttccga
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Cys Gly Gln Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Phe
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Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala

Ala	Glv	Leu	Lvs	Asp	Cys	Thr	Met	Leu	Val	Cys	Gly	Asp	Asp	Leu	Val
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	4 .	•		
cctgactaat tcaaaggggc agaactgtgg	ctatcgccga	tgccgcgcaa	gcggcgtgct	180
gacgaccage tgcggtaata cccttacatg	ttacctaaag	gcctctgcag	cctgtcgagc	240

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Glu Arg Leu Tyr Ile Gly Gly Pro Leu Thr Asn Ser Lys Gly Gln Asn 35

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Thr Thr Ser Cys

Gly 65	Asn	Thr	Leu		Cys 70	Tyr	Leu	Lys	Ala	Ser 75	Ala	Ala	Cys	Arg	Ala 80	
Ala	Lys	Leu	Gln	Asp 85	Cys	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
Val	Ile	Cys	Glu 100	Ser	Ala	Gly	Thr	Gln 105	Glu	Asp	Ala	Ala	Ser 110	Leu	Arg	
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cat	gata	aac	agca	aagg	gc a	atco	tgcg	g tt	acag	gcgt	. tgc	cgcg	caa	gcgg	tgtttt	180
															taaagc	240
															ctcgga	300
		gggt			,											340
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Al	a Cy	s Se	r Leu 20	ı Pro	Gl:	n Gl	u Ala	a Ar	g _. Th:	r Va	1 11	e Hi	s Se:	r Le	u Thr	· T
Gl	u Ar	g Le	u Tyi	r Val	. Gl	y Gl	y Pr 40		t Il	e As	n Se	r Ly 45	s Gl	y Gl:	n Ser	

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Met 50 55 60

Gly Asn 65	Thr	Met	Thr	Cys 70	Tyr	Ile	Lys	Ala	Leu 75	Ala	Ala	Cys	Lys	Ala 80	
Ala Gly	Ile	Val	Asp 85	Pro	Val	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
Val Ile	Ser	Glu 100	Ser	Gln	Gly	Asn	Glu 105	Glu	Asp.	Glu	Arg	Asn 110	Leu	Arg	
Ala			· •												
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gcctaco	cagc	tttg	gcaa	ca c	aatc	actt	g tt	acat	caag	gcc	acaa	cgg	ccgc	gaaggc	240
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Cys Cy	s Ası	n Lev 20	ı Glu	ı Pro	o Gli	ı Ala	a Arg 25	д Гу	s Va	1 I1	e Sei	Se:	r Lei	ı Thr	
Glu Ar	g Le 35	u Tyi	c Cys	s Gl	y Gl	y Pro	o Me	t Ph	e As	n Se	r Ly: 45	s Gl	y Ala	a Gln	
Cys Gl 50		r Arg	g Ar	g Cy:	s Arg	g Al	a Se	r Gl	y Va	1 Le [.]	u Pr	o Th	r Se	r Phe	

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cccatatat	cgaacaagct	caggtaatag	ctcaccagtt	caaggaaaaa	gtccttggat	540
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Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro Pro Ser 35 40 45

Trp Asp Glu Met Trp Lys Cys Leu Val Arg Leu Lys Pro Thr Leu His
50 55 60

Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile 65 70 75 80

Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser Ala 85 90 95

Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu Gly Gly Val Leu 100 105 110

Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys Val Val Ile Val 115 120 125

Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val Pro Asp Lys Glu 130 135 140

Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala 145 150 155 160

Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Glu Lys 165 170 175

Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Ala Val Ile Glu 180 185 190

Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala Phe Trp His Lys 195 200 205

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Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr 35 40 45

Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala
50 55 60

Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu 65 70 75 80

Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Gly Arg Ala 85 90 95

Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His
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Ser Lys Lys Lys Cys Asp Glu Léu Ala Lys Gln Leu Thr Ser Leu Gly 115 120 125

Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala Val Ile Pro 130 135 140 Thr Ala Gly Asp Val Val Cys Ser Thr Asp Ala Leu Met Thr Gly
145 150 155 160

Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser Ala Val Thr 165 170 175

Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Glu Thr Thr 180 185 190

Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg Gly Arg Thr 195 200 205

Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala Gly Glu Arg 210 215 220

Pro Ser Asp Met Phe Asp Ser Val Val Leu Cys Glu Cys Tyr Asp Ala 225 230 235 240

Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr Val Arg Leu 245 250 255

Arg Ala Tyr Ile Asn Thr Pro Gly Leu Pro Val Cys Gln Asp His Leu 260 265 270

Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile Asp Ala His 275 280 285

Met Leu Ser Gln Thr Lys Gln Gly Glu Asn Phe Pro Tyr Leu Val 290 295 300

Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro Pro Pro Ser 305 310 315 320

Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro Thr Leu Thr 325 330 335

Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile 340 345 350

Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser Ala 355 360 365

Asp Leu Glu Val Ile Thr Ser Thr Trp Val Leu Val Gly Gly Val Val 370 380

Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val Ala Ile Val 385 390 395 400

Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu 405 410 415

Ala Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser Ala Ser Leu 420 425 430

Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe Lys Glu Lys 435 440 445

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Glu Phe Asp Glu

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Thr Lys Gln Ala
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Ala Phe Asp Glu
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Ser Lys Gln Ala 20

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Leu Lys Ser Lys 20

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Thr Arg Gln Ala 20

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Phe Lys Glu Lys 20